SEQUENCE LISTING

```
<110> Stapleton, Jack
      Xiang, Jinhua
      George, Sarah
      University of Iowa Research Foundation
<120> GB VIRUS C AND METHODS OF TREATING VIRAL INFECTIONS
<130> 875.076US1
<150> US 60/421,408
<151> 2002-10-24
<160> 44
<170> FastSEQ for Windows Version 4.0
<210> 1
<211> 9395
<212> DNA
<213> Hepatitis G virus
<400> 1
tgacgtgggg gggttgatcc ccccccccg gcactgggtg caagccccat aaaccgacgc
                                                                        60
ctatctaagt agacgcaatg actcggcgc gactcggcga ccggccaaaa ggtggtggat
                                                                       120
gggtggtgac agggttggta ggtcgtaaat cccggtcatc ctggtagcca ctataggtgg
                                                                       180
gtcttaagag aaggtcaaga ctcctcttgt gcctgcggcg agaccgcgca cggtccacag
                                                                       240
gtgctggccc taccggtgtg aataagggcc cgacgtcagg ctcgtcgtta aaccgagccc
                                                                       300
gtcacccacc tgggcaaacg acgcccacgt acggtccacg tcgcccttca atgtctctct
                                                                       360
tgaccaatag gtttatccgg cgagttgaca aggaccagtg ggggccgggg gttatgggga
                                                                       420
aggaccccaa accetgeeet teeeggtggg eegggaaatg catggggeea eecageteeg
                                                                       480
cggcggcctg cagccggggt agcccaagaa tccttcgggt gagggcgggt ggcatttctc
                                                                       540
ttttctatac catcatggca gtccttctgc tccttctcgt ggttgaggcc ggggccattc
                                                                       600
tggccccggc cacccacgct tgtcgagcga atgggcaata tttcctcaca aattgctgtg
                                                                       660
ccccggaaga catcgggttc tgcctggaag gcggatgcct ggtggccctg gggtgcacgg
                                                                       720
tttgcaccga ccgttgctgg ccactgtatc aggcgggttt ggctgtgcgg cctqqcaaqt
                                                                       780
ccgcggccca gctcgttggg gaactgggga gcctgtacgg gcccttgtcg gtctcqqctt
                                                                       840
acgtagccgg gatcctgggt ctgggcgagg tttactccgg ggtcctgaca gttggtgttg
                                                                       900
cgttgaggcg ccgggtctac ctgatgccca acctgaagtg tgcagtagaa tgtgacgtta
                                                                       960
agtggggaag tgagttttgg agatggactg agcagttggc ctccaattac tggattttgg
                                                                      1020
aatacctttg gaaagtccca tttgaatttt ggagaggagt gatgagcctg acccctctgt
                                                                      1080
tggtttgggt ggccgcattg cttttgctgg agcaacggat tgtcatggtt ttcctgctgg
                                                                      1140
tgacgatggc ggggatgttg caaggcgccc ccgcctccgt tttggggtcc cgccctttg
                                                                      1200
actacgggtt gaagtggcag tcatgctcct gcagggctaa cgggtcgcgt attcccactg
                                                                      1260
gggagagggt gtgggatcga gggaatgtca cgctcttgtg tgactgcccc aacggcccct
                                                                      1320
gggtttgggt cccggccttt tgccaggcgg ttgggtgggg cgaccccatc acccattgga
                                                                      1380
gccacggaca aaaccagtgg cccctatcat gcccccaata tgtctatggg tctgtgtccg
                                                                      1440
taacgtgcgt gtggggttcc gtgtcttggt ttgcctcgac cggcggtcgt gattcgaaga
                                                                      1500
tegatgtgtg gagtttggtg ceggttggat etgecagetg caccatagee getetagggt
                                                                      1560
categgateg egacaeggtg gttgagetet eegagtgggg agteeegtge gtaaegtgta
                                                                      1620
ttctggaccg tcggcctgct tcatgtggca cctgtgtgcg ggactgctgg cccgaaaccg
                                                                      1680
ggtcggttag attccctttc catcggtgcg gcacggggcc tcggctgaca aaggacttgg
                                                                      1740
aagetgtgee ettegteaae aggaeaaete eetteaeeat aaggggeeee etgggeaaee
                                                                      1800
aggggagagg caacceggtg eggtegeece tgggttttgg gteetacace atqaccaaqa
                                                                      1860
tccgggattc cctgcatttg gtgaaatgtc ccacaccagc catagagcct ccgactggaa
                                                                      1920
cgttcgggtt cttccccgga gtcccgccca ttaacaactg catgccgcta ggcacggaag
                                                                      1980
tgtctgaggc attgggcgga gctgggctta cgggggggtt ctacgagcct ctggttcgca
                                                                      2040
ggtgttcgga gctgatggga cgccgaaatc cggtttgccc ggggtacgca tggctgtcct
                                                                      2100
ctggtagacc tgacgggttc atacacgtcc aggggcacct gcaggaggtg gatgcgggca
                                                                      2160
acttcatccc tcctccacgc tggttgctct tggattttgt atttgtcctg ctctatctga
                                                                      2220
tgaagctggc tgaggcacgg ttggtcccgt tgatcttgct tctgctgtgg tggtgggtga
                                                                      2280
```

accagttggc ggttctagga ctgccggctg tggacgctgc cgtggcgggt gaagtttttg

2340

```
egggeeetge ettgteatgg tgtttgggee tteecactgt eagtatgata etaggtetag
                                                                      2400
caaacctggt gttgtacttt cggtggatgg gccctcagcg cctcatgttc ctcgtgttgt
                                                                      2460
ggaagctcgc tcggggagct ttcccgctgg cacttttgat ggggatttcg gcgacccgcg
                                                                      2520
ggcgcacctc tgtgctcggg gccgagttct gcttcgatgt cacattcgag gtggacactt
                                                                      2580
cggtgttggg ctgggtggtg gccagcgtgg tggcttgggc catagcgctc ctgagctcaa
                                                                      2640
tgagcgcagg ggggtggaag cacaaggccg tgatctatag gacgtggtgt aaagggtacc
                                                                      2700
aggetgtgcg ccagagggtg gtgcggagcc ccctcgggga ggggcgtcct accaagcttc
                                                                      2760
tgacgttcgc ctggtgcttg gcctcataca tctggccgga tgctgtgatg atggtggtgg
                                                                      2820
tggccttggt cctcctcttc ggcctgttcg acgcactgga ctgggccctg gaggagctcc
                                                                      2880
tggtctcccg gccctcgtta cggcgactgg cacgggtggt tgagtgctgt gtgatggcgg
                                                                      2940
gcgagaaggc caccaccatc cgactggtct ccaagatgtg cgcaagaggg gcctacctgt
                                                                      3000
ttgaccacat gggctctttc tcgcgcgctg tcaaggagcg cttgttggaa tgggacgcgg
                                                                      3060
ctttggagcc cttgtcattc actaggacgg actgtcgcat catcagagat gccgcgagga
                                                                      3120
ccctgtcctg cggacagtgc gtcatgggtt tacccgtggt agcacggcgc ggtgatgagg
                                                                      3180
ttctcatcgg cgtctttcag gatgtgaatc atttgcctcc cgggtttgtc ccgactgcac
                                                                      3240
cagttgtcat ccgtcggtgc ggaaagggct tcctgggggt cacgaaggca gccttgacag
                                                                      3300
gtagggatcc tgacttacat ccagggaacg tcatggtgtt ggggacggct acgtcacgaa
                                                                      3360
gcatgggcac atgtctgaat ggcctgctgt tcacaacttt ccatggggct tcatcccgaa
                                                                      3420
ccatcgccac gcccgtgggg gcccttaatc ccaggtggtg gtcagccagt gatgacgtca
                                                                      3480
cggtgtaccc gcttccagat ggggcaactt cgttgacgcc ctgcacttgc caggcggagt
                                                                      3540
cctgttgggt tattagatcc gacggggctt tgtgccatgg cttgagcaag ggggacaagg
                                                                      3600
ttgagctgga tgtggccatg gaggtctctg acttccgtgg ttcgtctggt tcaccggtcc
                                                                      3660
tttgcgacaa agggcacgca gtaagaatgc tcgtgtcagt gctccactct ggcggcaggg
                                                                      3720
ttactgcggc gcgattcact aggccgtgga ctcaagtacc aacagatgcc aagactacca
                                                                      3780
cagaaccccc tccggtgccg gcaaaaggag ttttcaagga ggccccgttg tttatgccta
                                                                      3840
cggggggggg aaagagcacc cgcgtaccgt tggagtacgg caacatgggc cacaaggtct
                                                                      3900
tgatcttgaa cccgtcggta gctaccgtga gggccatggg cccatacatg gagcggctgg
                                                                      3960
cggggaaaca ccccagtatt tactgtggcc atgacaccac tgctttcaca aggatcactg
                                                                      4020
actogoccot tacgtattoc acttacggaa ggtttttggc caaccotagg cagatgotga
                                                                      4080
ggggtgtgtc ggtggtcatt tgtgacgagt gccacagtca tgactcaact gtgttqttqq
                                                                      4140
gcattgggcg tgtcagggag ctggcgcgag gatgtggagt gcaattggtg ctctacgcca
                                                                      4200
etgecaceee teeeggatee eegatgaeee ageaceeate aateattgag acaaaaetgg
                                                                      4260
acgtgggaga gatccccttc tatgggcatg gcatacctct tgagcggatg cggaccggaa
                                                                      4320
ggcatctcgt attctgccac tccaaggctg agtgcgagcg cctggcgggc cagttttcgg
                                                                      4380
ctaggggggt aaatgccatc gcctattaca gggggaaaga cagttctatc atcaaagatg
                                                                      4440
gagacctggt ggtgtgtgct acagacgcac tatccactgg gtacactggg aacttcgatt
                                                                      4500
ctgtcaccga ttgtgggtta gtggtggagg aggtcgtcga ggtgaccctt gatcccacca
                                                                      4560
ttaccatctc cctgcgcacg gtgcccgcgt cggctgaact gtcgatgcag cggcgaggac
                                                                      4620
gcacgggtag gggcaggtct gggcgctact actacgcggg ggtcggcaag gcccctgctg
                                                                      4680
gtgtggtgcg ctcaggtcct gtctggtcgg cggtggaagc cggtgtgacc tggtacggaa
                                                                      4740
tggaacctga cctgacagca aacctactga gactttacga caactgccct tacaccgcag
                                                                      4800
ccgtcgcagc tgacattggg gaagccgcgg tgttcttttc ggggcttgcc ccgttgagga
                                                                      4860
tgcatcccga tgttagctgg gcaaaagttc gcggcgtcaa ctggcccttc ctggtgggtg
                                                                      4920
ttcagcggac catgtgccgg gaaacactgt ctcccggccc atcggatgac ccccagtggg
                                                                      4980
caggtotgaa gggcccgaat cotgtoccac tootgotgag gtggggcaat gatttaccat
                                                                      5040
ctaaagtggc cggccatcac atcgtggacg acctggtccg taggctcggg gtggcggagg
                                                                      5100
gttacgtccg ctgcgatgcg ggacccatct tgatggtggg cctcgctatt gcggggggca
                                                                      5160
tgatctatgc gtcatacacc gggtctctcg tggtggttac agactgggat gtgaaggggg
                                                                      5220
gtggcagccc cctttatcgg catggagacc aggccacgcc ccagccggtt gtgcaggtcc
                                                                      5280
ecceggtaga ceateggeeg gggggagagt etgegeeate ggatgeeaac acagtgaeag
                                                                      5340
atgcggtggc ggccatccag gtggattgcg attggtcagt catgaccctg tcgatcgggg
                                                                      5400
aagtgctgtc cttggcccag gctaagacgg ccgaggccta cgcagctacc accaagtggc
                                                                      5460
ttgctggctg ctacacgggg acgcgggccg tccccactgt ttcaattgtt gacaagctct
                                                                      5520
tegeeggggg etgggeggeg gtggtaggee attgceaeag tgtaataget geggeagtgg
                                                                      5580
eggeetatgg ggettetagg ageetteeat tggetgetge egetteetae eteatggggt
                                                                      5640
tgggcgtcgg aggcaacgcg caaacccgct tagcctccgc tctcctacta ggggccgctg
                                                                      5700
ggaccgctct gggcacgcct gtcgtggggt taaccatggc gggcgcgttc atgggaagtg
                                                                      5760
ctagegtete ecceteettg gteaceattt taetggggge egtgggggge tgggagggeg
                                                                      5820
tggtgaatgc ggctagcctt gtcttcgact ttatggcggg gaaactatca tcagaagatc
                                                                      5880
tgtggtatgc catcccagtg ctaaccagtc cgggggcagg acttgcgggg atcgccctcg
                                                                      5940
ggttggtgtt gtactcagct aacaactctg gcactaccac ttggttgaac cgtctgctga
                                                                      6000
ctacattgcc aaggtcctca tgcatccctg acagttactt tcagcaggcc gattactgtg
                                                                      6060
```

acaaggtete agetgtgete egacgettga geeteacteg cacegtggtt geeetggtea 6120 acagggagcc taaggtggat gaggttcagg tggggtacgt ctgggacttg tgggagtgga 6180 teatgegtea agtgegeatg gtgatggeea gaetteggge cetetgeece gtggtgteat 6240 taccettatg gcactgcggg gaggggtggt ccggagaatg gttgttggac ggccatgttg 6300 agagtcgttg tctttgtggt tgcgtgatca ccggtgatgt tttgaatggg caactcaaag 6360 atccagttta ctctaccaag ctgtgcaggc attattggat ggggacagtc cctgtgaaca 6420 tgctgggcta tggcgagacg tcgcctttgc tcgcctcaga caccccgaag gtggtaccat 6480 tegggaegte tgggtggget gaggtggtgg tgaccectae ceaegttgtg ateaggegaa 6540 catecgccta caaactgctg cgccagcaaa tcctgtcggc tgctgttgct gagccctatt 6600 acgtcgacgg cataccggtc tcatgggacg cggacgcgcg agcgcctgcc atggtctatg 6660 gecetgggea aagtgteace attgaegggg aaegetacae eetteegeat caaetgegge 6720 ttaggaatgt ggcgccctct gaggtgtcat ccgaggtgtc cattgacatt gggacggaga 6780 ctgaagactc agaactgact gaggccgacc tgccgccggc ggctgcagcc cttcaggcta 6840 tcgagaatgc tgcgagaatt cttgaacctc acatagatgt catcatggaa gattgcagta 6900 caccetetet ttgtgggagt ageegagaga tgeetgtgtg gggagaagae atacceegea 6960 ctccatcgcc agcacttatc tcggttactg agagcagccc agatgagaag accccqtcqq 7020 tgtcttcctc gcaggaggat accccgtctt ctgactcatt cgaggtcatc caaqaqtccq 7080 agacagcega aggggaggaa agegtettea aegtggetet tteegtaeta aaaqeettqt 7140 ttccacagag cgatgccaca agaaagctta ccgttaagat gtcatgctgt gttgagaaga 7200 gcgtaacacg cttcttttca ttgggattga cggtcgctga cgtggcaagc ctqtqtqaqa 7260 tggaaatcca gaaccataca gcctattgtg acaaggtgcg cactccgctt gaattgcagg 7320 ttgggtgctt ggtgggcaat gaacttacct ttgaatgtga caagtgtgag gctaggcaag 7380 agacettgge tteettetet tacatttggt etggggtgee aetgaegagg geeaeteegg 7440 ccaagccccc tgtggtgagg ccggttggct ccttgctggt ggccgacacc accaaggtgt 7500 atgtcaccaa cccggacaat gttgggagaa gagttgacaa ggttaccttc tggcgtqccc 7560 ctagggttca tgacaaattc ctcgtggact ccatagagcg cgctaagagg gcagctcaag 7620 7680 cctgcctaag catgggttac acttatgagg aggcaataag gactgtaagg ccacatgctg ccatgggctg gggatctaag gtgtcggtca aggacctcgc cacccctgcg gggaagatqq 7740 etgtecatga ceggetecag gagataettg aagggaegee agteceettt actettaetg 7800 tgaaaaagga agtgttcttc aaagaccgaa aggaagagaa ggccccccgc ctcattgtgt 7860 tecececet ggaetteegg atagetgaaa agettattet gggagaeeet ggaegggtag 7920 7980 ccaaggcggt gttggggggg gcctacgcct tccagtacac cccaaatcag cgaattaggg agatgeteaa aetgtgggaa teaaagaaga caecatgege eatetgtgtg gaegeeacat 8040 gcttcgacag tagcataact gaagaggacg tggcgctgga gacagagctt tatgccctgg 8100 cttcagacca tccagaatgg gtgcgtgccc tggggaaata ctatgcctct ggcacaatgg 8160 taacccccga gggggtgcca gtgggtgaga ggtattgtag atcctcaggg gtcttgacca 8220 ccagtgcgag caactgcttg acttgctata tcaaggtgaa agccgcctgt gagagggtgg 8280 ggctgaaaaa tgtctcgctc ctcatcgctg gcgatgactg tttgatcata tgcgaacggc 8340 ctgtgtgcga tcctagcgac gctttgggca gagccctggc gagctacggg tacgcatgcg 8400 ageettegta teatgeatea etggacaegg ecceettetg etceaettgg etagetgagt 8460 gcaatgcaga tgggaaacgc catttcttcc tgaccacgga ctttcggagg cccctcgctc 8520 gcatgtcgag cgagtacagt gacccaatgg cttcggccat cggttacatc ctcctatacc 8580 cttggcatcc tatcacacgg tgggtcatca tccctcacgt gctcacctgc gcgtttaggg 8640 gtggtggcac accgtctgat cctgtgtggt gccaggtaca tggtaattac tacaagtttc 8700 cactggacaa actgcctaac atcatcgtgg ccctccacgg accagcagcg ttgagggtta 8760 ccgcagacac aactaagaca aaaatggagg ctggcaaggt gctqagcgac ctcaagctcc 8820 ctggcctagc agtccaccgg aagaaggccg gggcattgcg aacgcgtatg ctccggtcgc 8880 gcggttgggc tgagttggct agggggctgt tgtggcgtcc aggcctgcgg cttcccctc 8940 eggagattge tggtateece gggggtttee ceettteece eccetatatg ggggttgt 9000 atcaattgga tttcacaagc cagaggagtc gctggcggtg gttggggttc ttagccctgc 9060 tcatcgtagc cctcttcggg tgaactaaat tcatctgttg cggcaaggtc cggtgactga 9120 teateactgg aggaggttee egeceteece geeceagggg teteceeget gggtaaaaag 9180 ggcccggcct tgggaggcat ggtggttact aaccccctgg cagggtcaaa gcctgatggt 9240 gctaatgcac tgccacttcg gtggcgggtc gctaccttat agcgtaatcc gtgactacgg 9300 gctgctcgca gagccctccc cggatggggc acagtgcact gtgatctgaa ggggtgcacc 9360 ccggtaagag ctcggcccaa aggccgggtt ctact 9395

```
<212> PRT
<213> Hepatitis G virus
<400> 2
Met Ser Leu Leu Thr Asn Arg Phe Ile Arg Arg Val Asp Lys Asp Gln
                                    10
Trp Gly Pro Gly Val Met Gly Lys Asp Pro Lys Pro Cys Pro Ser Arg
                                25
Trp Ala Gly Lys Cys Met Gly Pro Pro Ser Ser Ala Ala Ala Cys Ser
Arg Gly Ser Pro Arg Ile Leu Arg Val Arg Ala Gly Gly Ile Ser Leu
Phe Tyr Thr Ile Met Ala Val Leu Leu Leu Leu Val Val Glu Ala
                                         75
Gly Ala Ile Leu Ala Pro Ala Thr His Ala Cys Arg Ala Asn Gly Gln
Tyr Phe Leu Thr Asn Cys Cys Ala Pro Glu Asp Ile Gly Phe Cys Leu
                                105
Glu Gly Gly Cys Leu Val Ala Leu Gly Cys Thr Val Cys Thr Asp Arg
                            120
                                                 125
Cys Trp Pro Leu Tyr Gln Ala Gly Leu Ala Val Arg Pro Gly Lys Ser
    130
Ala Ala Gln Leu Val Gly Glu Leu Gly Ser Leu Tyr Gly Pro Leu Ser
                    150
Val Ser Ala Tyr Val Ala Gly Ile Leu Gly Leu Gly Glu Val Tyr Ser
                                    170
Gly Val Leu Thr Val Gly Val Ala Leu Arg Arg Arg Val Tyr Leu Met
            180
                                                     190
Pro Asn Leu Lys Cys Ala Val Glu Cys Asp Val Lys Trp Gly Ser Glu
                            200
                                                 205
Phe Trp Arg Trp Thr Glu Gln Leu Ala Ser Asn Tyr Trp Ile Leu Glu
                        215
                                            220
Tyr Leu Trp Lys Val Pro Phe Glu Phe Trp Arg Gly Val Met Ser Leu
                    230
                                        235
Thr Pro Leu Leu Val Trp Val Ala Ala Leu Leu Leu Leu Glu Gln Arg
                245
                                    250
Ile Val Met Val Phe Leu Leu Val Thr Met Ala Gly Met Leu Gln Gly
                                265
                                                     270
Ala Pro Ala Ser Val Leu Gly Ser Arg Pro Phe Asp Tyr Gly Leu Lys
                            280
                                                285
Trp Gln Ser Cys Ser Cys Arg Ala Asn Gly Ser Arg Ile Pro Thr Gly
                        295
                                            300
Glu Arg Val Trp Asp Arg Gly Asn Val Thr Leu Leu Cys Asp Cys Pro
                    310
                                        315
Asn Gly Pro Trp Val Trp Val Pro Ala Phe Cys Gln Ala Val Gly Trp
                                    330
Gly Asp Pro Ile Thr His Trp Ser His Gly Gln Asn Gln Trp Pro Leu
                                345
                                                    350
Ser Cys Pro Gln Tyr Val Tyr Gly Ser Val Ser Val Thr Cys Val Trp
                            360
Gly Ser Val Ser Trp Phe Ala Ser Thr Gly Gly Arg Asp Ser Lys Ile
                        375
Asp Val Trp Ser Leu Val Pro Val Gly Ser Ala Ser Cys Thr Ile Ala
                    390
Ala Leu Gly Ser Ser Asp Arg Asp Thr Val Val Glu Leu Ser Glu Trp
                405
                                    410
Gly Val Pro Cys Val Thr Cys Ile Leu Asp Arg Arg Pro Ala Ser Cys
           420
                                425
Gly Thr Cys Val Arg Asp Cys Trp Pro Glu Thr Gly Ser Val Arg Phe
        435
                            440
```

<210> 2 <211> 2910

Pro Phe His Arg Cys Gly Thr Gly Pro Arg Leu Thr Lys Asp Leu Glu Ala Val Pro Phe Val Asn Arg Thr Thr Pro Phe Thr Ile Arg Gly Pro Leu Gly Asn Gln Gly Arg Gly Asn Pro Val Arg Ser Pro Leu Gly Phe Gly Ser Tyr Thr Met Thr Lys Ile Arg Asp Ser Leu His Leu Val Lys Cys Pro Thr Pro Ala Ile Glu Pro Pro Thr Gly Thr Phe Gly Phe Phe Pro Gly Val Pro Pro Ile Asn Asn Cys Met Pro Leu Gly Thr Glu Val Ser Glu Ala Leu Gly Gly Ala Gly Leu Thr Gly Gly Phe Tyr Glu Pro Leu Val Arg Arg Cys Ser Glu Leu Met Gly Arg Arg Asn Pro Val Cys Pro Gly Tyr Ala Trp Leu Ser Ser Gly Arg Pro Asp Gly Phe Ile His Val Gln Gly His Leu Gln Glu Val Asp Ala Gly Asn Phe Ile Pro Pro Pro Arg Trp Leu Leu Asp Phe Val Phe Val Leu Leu Tyr Leu Met Lys Leu Ala Glu Ala Arg Leu Val Pro Leu Ile Leu Leu Leu Trp Trp Trp Val Asn Gln Leu Ala Val Leu Gly Leu Pro Ala Val Asp Ala Ala Val Ala Gly Glu Val Phe Ala Gly Pro Ala Leu Ser Trp Cys Leu Gly Leu Pro Thr Val Ser Met Ile Leu Gly Leu Ala Asn Leu Val Leu Tyr Phe Arg Trp Met Gly Pro Gln Arg Leu Met Phe Leu Val Leu Trp Lys Leu Ala Arg Gly Ala Phe Pro Leu Ala Leu Leu Met Gly Ile Ser Ala Thr Arg Gly Arg Thr Ser Val Leu Gly Ala Glu Phe Cys Phe Asp Val Thr Phe Glu Val Asp Thr Ser Val Leu Gly Trp Val Val Ala Ser Val Val Ala Trp Ala Ile Ala Leu Leu Ser Ser Met Ser Ala Gly Gly Trp Lys His Lys Ala Val Ile Tyr Arg Thr Trp Cys Lys Gly Tyr Gln Ala Val Arg Gln Arg Val Val Arg Ser Pro Leu Gly Glu Gly Arg Pro Thr Lys Leu Leu Thr Phe Ala Trp Cys Leu Ala Ser Tyr Ile Trp Pro Asp Ala Val Met Met Val Val Val Leu Val Leu Leu Phe Gly Leu Phe Asp Ala Leu Asp Trp Ala Leu Glu Glu Leu Leu Val Ser Arg Pro Ser Leu Arg Arg Leu Ala Arg Val Val Glu Cys Cys Val Met Ala Gly Glu Lys Ala Thr Thr Ile Arg Leu Val Ser Lys Met Cys Ala Arg Gly Ala Tyr Leu Phe Asp His Met Gly Ser Phe Ser Arg Ala Val Lys Glu Arg Leu Leu Glu Trp Asp Ala Ala Leu Glu Pro Leu Ser Phe Thr Arg Thr Asp Cys Arg Ile Ile Arg Asp Ala Ala Arg Thr Leu Ser Cys Gly Gln Cys Val Met Gly Leu Pro Val Val Ala Arg Arg Gly Asp Glu Val

```
Leu Ile Gly Val Phe Gln Asp Val Asn His Leu Pro Pro Gly Phe Val
945
                   950
                                       955
Pro Thr Ala Pro Val Val Ile Arg Arg Cys Gly Lys Gly Phe Leu Gly
                965
                                   970
Val Thr Lys Ala Ala Leu Thr Gly Arg Asp Pro Asp Leu His Pro Gly
                              985
Asn Val Met Val Leu Gly Thr Ala Thr Ser Arg Ser Met Gly Thr Cys
                           1000
                                          1005
Leu Asn Gly Leu Leu Phe Thr Thr Phe His Gly Ala Ser Ser Arg Thr
                       1015
                                         1020
Ile Ala Thr Pro Val Gly Ala Leu Asn Pro Arg Trp Trp Ser Ala Ser
                   1030
                                      1035
Asp Asp Val Thr Val Tyr Pro Leu Pro Asp Gly Ala Thr Ser Leu Thr
               1045
                                  1050
Pro Cys Thr Cys Gln Ala Glu Ser Cys Trp Val Ile Arg Ser Asp Gly
           1060
                              1065
Ala Leu Cys His Gly Leu Ser Lys Gly Asp Lys Val Glu Leu Asp Val
        1075
                          1080
Ala Met Glu Val Ser Asp Phe Arg Gly Ser Ser Gly Ser Pro Val Leu
                       1095
                                          1100
Cys Asp Lys Gly His Ala Val Arg Met Leu Val Ser Val Leu His Ser
                   1110
                                      1115
Gly Gly Arg Val Thr Ala Ala Arg Phe Thr Arg Pro Trp Thr Gln Val
               1125
                                  1130
Pro Thr Asp Ala Lys Thr Thr Glu Pro Pro Pro Val Pro Ala Lys
           1140
                               1145
                                                  1150
Gly Val Phe Lys Glu Ala Pro Leu Phe Met Pro Thr Gly Ala Gly Lys
       1155
                           1160
                                              1165
Ser Thr Arg Val Pro Leu Glu Tyr Gly Asn Met Gly His Lys Val Leu
                       1175
                                          1180
Ile Leu Asn Pro Ser Val Ala Thr Val Arg Ala Met Gly Pro Tyr Met
                   1190
                                     1195
Glu Arg Leu Ala Gly Lys His Pro Ser Ile Tyr Cys Gly His Asp Thr
               1205
                                  1210
                                                     1215
Thr Ala Phe Thr Arg Ile Thr Asp Ser Pro Leu Thr Tyr Ser Thr Tyr
           1220
                             1225
                                                  1230
Gly Arg Phe Leu Ala Asn Pro Arg Gln Met Leu Arg Gly Val Ser Val
       1235
                          1240
                                             1245
Val Ile Cys Asp Glu Cys His Ser His Asp Ser Thr Val Leu Leu Gly
                      1255
                                          1260
Ile Gly Arg Val Arg Glu Leu Ala Arg Gly Cys Gly Val Gln Leu Val
                   1270
                                     1275
Leu Tyr Ala Thr Ala Thr Pro Pro Gly Ser Pro Met Thr Gln His Pro
               1285
                                  1290
                                                      1295
Ser Ile Ile Glu Thr Lys Leu Asp Val Gly Glu Ile Pro Phe Tyr Gly
           1300
                              1305
                                                  1310
His Gly Ile Pro Leu Glu Arg Met Arg Thr Gly Arg His Leu Val Phe
                          1320
                                              1325
Cys His Ser Lys Ala Glu Cys Glu Arg Leu Ala Gly Gln Phe Ser Ala
                       1335
                                          1340
Arg Gly Val Asn Ala Ile Ala Tyr Tyr Arg Gly Lys Asp Ser Ser Ile
                   1350
                                      1355
Ile Lys Asp Gly Asp Leu Val Val Cys Ala Thr Asp Ala Leu Ser Thr
               1365
                                  1370
                                                      1375
Gly Tyr Thr Gly Asn Phe Asp Ser Val Thr Asp Cys Gly Leu Val Val
                              1385
                                                  1390
Glu Glu Val Val Glu Val Thr Leu Asp Pro Thr Ile Thr Ile Ser Leu
                           1400
                                              1405
Arg Thr Val Pro Ala Ser Ala Glu Leu Ser Met Gln Arg Arg Gly Arg
                       1415
                                          1420
Thr Gly Arg Gly Arg Ser Gly Arg Tyr Tyr Ala Gly Val Gly Lys
                   1430
                                      1435
```

Ala Pro Ala Gly Val Val Arg Ser Gly Pro Val Trp Ser Ala Val Glu Ala Gly Val Thr Trp Tyr Gly Met Glu Pro Asp Leu Thr Ala Asn Leu Leu Arg Leu Tyr Asp Asn Cys Pro Tyr Thr Ala Ala Val Ala Ala Asp Ile Gly Glu Ala Ala Val Phe Phe Ser Gly Leu Ala Pro Leu Arg Met His Pro Asp Val Ser Trp Ala Lys Val Arg Gly Val Asn Trp Pro Phe Leu Val Gly Val Gln Arg Thr Met Cys Arg Glu Thr Leu Ser Pro Gly Pro Ser Asp Asp Pro Gln Trp Ala Gly Leu Lys Gly Pro Asn Pro Val Pro Leu Leu Arg Trp Gly Asn Asp Leu Pro Ser Lys Val Ala Gly His His Ile Val Asp Asp Leu Val Arg Arg Leu Gly Val Ala Glu Gly Tyr Val Arg Cys Asp Ala Gly Pro Ile Leu Met Val Gly Leu Ala Ile Ala Gly Gly Met Ile Tyr Ala Ser Tyr Thr Gly Ser Leu Val Val Thr Asp Trp Asp Val Lys Gly Gly Ser Pro Leu Tyr Arg His Gly Asp Gln Ala Thr Pro Gln Pro Val Val Gln Val Pro Pro Val Asp His Arg Pro Gly Glu Ser Ala Pro Ser Asp Ala Asn Thr Val Thr Asp Ala Val Ala Ala Ile Gln Val Asp Cys Asp Trp Ser Val Met Thr Leu Ser Ile Gly Glu Val Leu Ser Leu Ala Gln Ala Lys Thr Ala Glu Ala Tyr Ala Ala Thr Thr Lys Trp Leu Ala Gly Cys Tyr Thr Gly Thr Arg Ala Val Pro Thr Val Ser Ile Val Asp Lys Leu Phe Ala Gly Gly Trp Ala Ala Val Val Gly His Cys His Ser Val Ile Ala Ala Ala Val Ala Ala Tyr Gly Ala Ser Arg Ser Pro Pro Leu Ala Ala Ala Ala Ser Tyr Leu Met Gly Leu Gly Val Gly Asn Ala Gln Thr Arg Leu Ala Ser Ala Leu Leu Gly Ala Ala Gly Thr Ala Leu Gly Thr Pro Val Val Gly Leu Thr Met Ala Gly Ala Phe Met Gly Ser Ala Ser Val Ser Pro Ser Leu Val Thr Ile Leu Leu Gly Ala Val Gly Gly Trp Glu Gly Val Val Asn Ala Ala Ser Leu Val Phe Asp Phe Met Ala Gly Lys Leu Ser Ser Glu Asp Leu Trp Tyr Ala Ile Pro Val Leu Thr Ser Pro Gly Ala Gly Leu Ala Gly Ile Ala Leu Gly Leu Val Leu Tyr Ser Ala Asn Asn Ser Gly Thr Thr Trp Leu Asn Arg Leu Leu Thr Thr Leu Pro Arg Ser Ser Cys Ile Pro Asp Ser Tyr Phe Gln Gln Ala Asp Tyr Cys Asp Lys Val Ser Ala Val Leu Arg Arg Leu Ser Leu Thr Arg Thr Val Val Ala Leu Val Asn Arg Glu Pro Lys Val Asp Glu Val Gln Val Gly Tyr

```
Val Trp Asp Leu Trp Glu Trp Ile Met Arg Gln Val Arg Met Val Met
          1940
                              1945
Ala Arg Leu Arg Ala Leu Cys Pro Val Val Ser Leu Pro Leu Trp His
       1955
                         1960
                                              1965
Cys Gly Glu Gly Trp Ser Gly Glu Trp Leu Leu Asp Gly His Val Glu
                      1975
                                          1980
Ser Arg Cys Leu Cys Gly Cys Val Ile Thr Gly Asp Val Leu Asn Gly
                1990
                                      1995
Gln Leu Lys Asp Pro Val Tyr Ser Thr Lys Leu Cys Arg His Tyr Trp
               2005
                                  2010
Met Gly Thr Val Pro Val Asn Met Leu Gly Tyr Gly Glu Thr Ser Pro
           2020
                              2025
                                                  2030
Leu Leu Ala Ser Asp Thr Pro Lys Val Val Pro Phe Gly Thr Ser Gly
                          2040
                                              2045
Trp Ala Glu Val Val Val Thr Pro Thr His Val Val Ile Arg Arg Thr
                      2055
                                          2060
Ser Ala Tyr Lys Leu Leu Arg Gln Gln Ile Leu Ser Ala Ala Val Ala
                  2070
                                      2075
Glu Pro Tyr Tyr Val Asp Gly Ile Pro Val Ser Trp Asp Ala Asp Ala
               2085
                                  2090
Arg Ala Pro Ala Met Val Tyr Gly Pro Gly Gln Ser Val Thr Ile Asp
                              2105
                                                  2110
Gly Glu Arg Tyr Thr Leu Pro His Gln Leu Arg Leu Arg Asn Val Ala
       2115
                          2120
                                              2125
Pro Ser Glu Val Ser Ser Glu Val Ser Ile Asp Ile Gly Thr Glu Thr
                      2135
                                          2140
Glu Asp Ser Glu Leu Thr Glu Ala Asp Leu Pro Pro Ala Ala Ala Ala
                   2150
                                      2155
Leu Gln Ala Ile Glu Asn Ala Ala Arg Ile Leu Glu Pro His Ile Asp
               2165
                                  2170
Val Ile Met Glu Asp Cys Ser Thr Pro Ser Leu Cys Gly Ser Ser Arg
           2180
                              2185
Glu Met Pro Val Trp Gly Glu Asp Ile Pro Arg Thr Pro Ser Pro Ala
                          2200
                                              2205
Leu Ile Ser Val Thr Glu Ser Ser Pro Asp Glu Lys Thr Pro Ser Val
                      2215
                                         2220
Ser Ser Ser Gln Glu Asp Thr Pro Ser Ser Asp Ser Phe Glu Val Ile
                  2230
                                      2235
Gln Glu Ser Glu Thr Ala Glu Gly Glu Glu Ser Val Phe Asn Val Ala
               2245
                                  2250
Leu Ser Val Leu Lys Ala Leu Phe Pro Gln Ser Asp Ala Thr Arg Lys
                              2265
Leu Thr Val Lys Met Ser Cys Cys Val Glu Lys Ser Val Thr Arg Phe
                          2280
Phe Ser Leu Gly Leu Thr Val Ala Asp Val Ala Ser Leu Cys Glu Met
                      2295
                                          2300
Glu Ile Gln Asn His Thr Ala Tyr Cys Asp Lys Val Arg Thr Pro Leu
                  2310
                                      2315
Glu Leu Gln Val Gly Cys Leu Val Gly Asn Glu Leu Thr Phe Glu Cys
               2325
                                  2330
Asp Lys Cys Glu Ala Arg Gln Glu Thr Leu Ala Ser Phe Ser Tyr Ile
           2340
                              2345
Trp Ser Gly Val Pro Leu Thr Arg Ala Thr Pro Ala Lys Pro Pro Val
                          2360
Val Arg Pro Val Gly Ser Leu Leu Val Ala Asp Thr Thr Lys Val Tyr
                     2375
                                          2380
Val Thr Asn Pro Asp Asn Val Gly Arg Arg Val Asp Lys Val Thr Phe
                 2390
                                      2395
Trp Arg Ala Pro Arg Val His Asp Lys Phe Leu Val Asp Ser Ile Glu
              2405
                                  2410
Arg Ala Lys Arg Ala Ala Gln Ala Cys Leu Ser Met Gly Tyr Thr Tyr
           2420
                              2425
```

Glu Glu Ala Ile Arg Thr Val Arg Pro His Ala Ala Met Gly Trp Gly Ser Lys Val Ser Val Lys Asp Leu Ala Thr Pro Ala Gly Lys Met Ala Val His Asp Arg Leu Gln Glu Ile Leu Glu Gly Thr Pro Val Pro Phe Thr Leu Thr Val Lys Lys Glu Val Phe Phe Lys Asp Arg Lys Glu Glu Lys Ala Pro Arg Leu Ile Val Phe Pro Pro Leu Asp Phe Arg Ile Ala Glu Lys Leu Ile Leu Gly Asp Pro Gly Arg Val Ala Lys Ala Val Leu Gly Gly Ala Tyr Ala Phe Gln Tyr Thr Pro Asn Gln Arg Ile Arg Glu Met Leu Lys Leu Trp Glu Ser Lys Lys Thr Pro Cys Ala Ile Cys Val Asp Ala Thr Cys Phe Asp Ser Ser Ile Thr Glu Glu Asp Val Ala Leu Glu Thr Glu Leu Tyr Ala Leu Ala Ser Asp His Pro Glu Trp Val Arg Ala Leu Gly Lys Tyr Tyr Ala Ser Gly Thr Met Val Thr Pro Glu Gly Val Pro Val Gly Glu Arg Tyr Cys Arg Ser Ser Gly Val Leu Thr Thr Ser Ala Ser Asn Cys Leu Thr Cys Tyr Ile Lys Val Lys Ala Ala Cys Glu Arg Val Gly Leu Lys Asn Val Ser Leu Leu Ile Ala Gly Asp Asp Cys Leu Ile Ile Cys Glu Arg Pro Val Cys Asp Pro Ser Asp Ala Leu Gly Arg Ala Leu Ala Ser Tyr Gly Tyr Ala Cys Glu Pro Ser Tyr His Ala Ser Leu Asp Thr Ala Pro Phe Cys Ser Thr Trp Leu Ala Glu Cys Asn Ala Asp Gly Lys Arg His Phe Phe Leu Thr Thr Asp Phe Arg Arg Pro Leu Ala Arg Met Ser Ser Glu Tyr Ser Asp Pro Met Ala Ser Ala Ile Gly Tyr Ile Leu Leu Tyr Pro Trp His Pro Ile Thr Arg Trp Val Ile Ile Pro His Val Leu Thr Cys Ala Phe Arg Gly Gly Thr Pro Ser Asp Pro Val Trp Cys Gln Val His Gly Asn Tyr Tyr Lys Phe Pro Leu Asp Lys Leu Pro Asn Ile Ile Val Ala Leu His Gly Pro Ala Ala Leu Arg Val Thr Ala Asp Thr Thr Lys Thr Lys Met Glu Ala Gly Lys Val Leu Ser Asp Leu Lys Leu Pro Gly Leu Ala Val His Arg Lys Lys Ala Gly Ala Leu Arg Thr Arg Met Leu Arg Ser Arg Gly Trp Ala Glu Leu Ala Arg Gly Leu Leu Trp Arg Pro Gly Leu Arg Leu Pro Pro Glu Ile Ala Gly Ile Pro Gly Gly Phe Pro Leu Ser Pro Pro Tyr Met Gly Val Val His Gln Leu Asp Phe Thr Ser Gln Arg Ser Arg Trp Arg Trp Leu Gly Phe Leu Ala Leu Leu Ile Val Ala Leu Phe Gly

```
<210> 3
<211> 36
<212> DNA
<213> Artificial Sequence
<220>
<223> A synthetic primer.
tcatggtggc gaataaaagc cccagaaacc gacgcc
                                                                          36
<210> 4
<211> 16
<212> DNA
<213> Artificial Sequence
<223> A synthetic primer.
<400> 4
tcatggtggc gaataa
                                                                          16
<210> 5
<211> 24
<212> DNA
<213> Artificial Sequence
<220>
<223> A synthetic primer.
<400> 5
tactgcartc ytccatgatg acat
                                                                          24
<210> 6
<211> 22
<212> DNA
<213> Artificial Sequence
<220>
<223> A synthetic primer.
<400> 6
atggtytayg gycctggvca aa
                                                                          22
<210> 7
<211> 23
<212> DNA
<213> Artificial Sequence
<220>
<223> A synthetic primer.
<400> 7
ttcaagaatc ctcgcagcat tct
                                                                          23
<210> 8
<211> 20
<212> DNA
<213> Artificial Sequence
<220>
<223> A synthetic primer.
```

<400> 8 ctggvcaaag ygtyaccatt	20
<210> 9 <211> 20 <212> DNA <213> Hepatitis G virus	
<400> 9 taccggtgtg aataagggcc	20
<210> 10 <211> 17 <212> DNA <213> Hepatitis G virus	
<400> 10 cgtcgtttgc ccaggtg	17
<210> 11 <211> 25 <212> DNA <213> Hepatitis G virus	
<400> 11 ctcgtcgtta aaccgagccc gtcac	25
<210> 12 <211> 24 <212> DNA <213> Artificial Sequence	
<220> <223> A synthetic primer.	
<400> 12 tactgcartc ytccatgatg acat	24
<210> 13 <211> 22 <212> DNA <213> Artificial Sequence	
<220> <223> A synthetic primer.	
<400> 13 atggtytayg gycctggvca aa	22
<210> 14 <211> 23 <212> DNA <213> Artificial Sequence	
<220> <223> A synthetic primer.	
<400> 14 ttcaagaatc ctcgcagcat tct	23
<210> 15	

```
<212> DNA
<213> Artificial Sequence
<220>
<223> A synthetic primer.
<400> 15
ctggvcaaag ygtyaccatt
                                                                         20
<210> 16
<211> 20
<212> DNA
<213> Hepatitis G virus
<400> 16
taccggtgtg aataagggcc
                                                                         20
<210> 17
<211> 17
<212> DNA
<213> Hepatitis G virus
<400> 17
cgtcgtttgc ccaggtg
                                                                         17
<210> 18
<211> 25
<212> DNA
<213> Hepatitis G virus
<400> 18
ctcgtcgtta aaccgagccc gtcac
                                                                        25
<210> 19
<211> 9309
<212> DNA
<213> Hepatitis G virus
<400> 19
aagccccaga aaccgacgcc tatctaagta gacgcaatga ctcggcgccg actcggcgac
                                                                        60
cggccaaaag gtggtggatg ggtgatgaca gggttggtag gtcgtaaatc ccggtcatcc
                                                                       120
tggtagccac tataggtggg tcttaagaga aggttaagat tcctcttgtg cctgcggcga
                                                                       180
gaccgcgcac ggtccacagg tgttggccct accggtgtga ataagggccc qacqtcaggc
                                                                       240
tegtegttaa acegageeeg ttacecacet gggeaaaega egeeeaegta eggteeaegt
                                                                       300
cgcccttcaa tgtctctctt gaccaatagg cttagccggc gagttgacaa ggaccagtgg
                                                                       360
gggccggggg ctatggagat ggactccaag tcctgccctt cccggtgggg cgggaaatgc
                                                                       420
atggggccac ccageteege ggeggeetge ageeggggta geecaagaac cettegggtg
                                                                       480
agggcgggtg gcatttttct tttctatacc atcatggcag tccttctgct ccttctcqtq
                                                                       540
gttgaggccg gggccatttt agccccggcc acccacgctt gtcgagcgaa tgggcaatac
                                                                       600
ttccttacaa attgttgcgc cccggaggac atcgggttct gcctggaagg tggatgcctg
                                                                       660
gtggccttgg gatgcacagt ttgcactgac caatgctggc cactgtatca ggcgggtttg
                                                                       720
gctgtgcggc ctggcaagtc cgcggcccaa ctggtggggg agctgggtag cctatacqqq
                                                                       780
cccttgtcgg tctcggccta cgtggctggg atcctgggcc tgggcgaggt ttactcgggt
                                                                       840
gtcctaacgg tgggcgtcgc gttgacgcgt cgggtctacc cggtgcccaa cctgacgtgt
                                                                       900
gcagtagagt gtgagcttaa gtgggaaagt gagttttgga gatggactga acagttggcc
                                                                       960
tccaattact ggattctgga atacctttgg aaggtcccat ttgacttctg gaggggcgtg
                                                                      1020
atgagectga eccetetgtt ggtetgtgtg geegetetge tgetgettga geaaaggatt
                                                                      1080
gttatggtct tcctgctggt gaccatggcc gggatgtcgc aaggcgcccc tgcttcggtt
                                                                      1140
ttggggtcac gcccctttga ctacgggttg acttggcaga cctgctcttg caaggcgaac
                                                                      1200
ggctcacgca ttccgactgg ggaaaaggtg tgggaccgcg ggaatgtcac gcttctgtgt
                                                                      1260
gattgcccca acggaccatg ggtgtggtta ccggctttct gccaagcagt tggctggggt
                                                                      1320
gatectatea eccattggag ecaeggacaa aateggtgge eccteteatg eccteagtat
                                                                      1380
```

```
gtctatgggt ctgtttcagt cacttgcgtg tggggctccg cgtcttggtt tgcctccact
                                                                      1440
ggtggacgcg attcgaagat cgatgtgtgg agtttggtgc cagttggttc tgccacttgc
                                                                      1500
accatagceg ctetegggte gteggacege gataeggtgg tggagetete egagtggggg
                                                                      1560
gtcccgtgcg tgacgtgcat tctggatcgt cggcctgcct cctgcggcac ctgtgtgagg
                                                                      1620
gactgctggc ccgagactgg gtcggtcaga ttcccattcc ataggtgtgg cacggggcct
                                                                      1680
cggctgacaa aggacttgga agctgtgccc ttcgtcaata ggacaactcc cttcaccatt
                                                                      1740
agggggcctc tgggcaacca gggccgaggc aacceggtgc ggtcgccctt gggttttggg
                                                                      1800
tectacacca tgaccaagat eegagacace etacatttgg tgaaatgtee cacaccagee
                                                                      1860
attgagcete ceaeegggae gtttgggtte tteeeeggga egeegeeet caacaaetge
                                                                      1920
atgettetag geaeggaagt gtetgaggea eteggegggg etggteteae ggggggttte
                                                                      1980
tatgaacccc tggtgcgcag gtgttcggag ctgatgggac gccgaaatcc ggtttgtccg
                                                                      2040
gggtttgcat ggctctcctc gggcaggcct gatgggttca tacatgtcca gggccacttg
                                                                      2100
                                                                      2160
caggaggtgg atgcaggcaa cttcattccg ccccgcgct ggctgctctt ggactttgta
tttgtcctgt tatacctgat gaagctggct gaggctcggt tggtcccgct gatcttgctt
                                                                      2220
ctgctgtggt ggtgggtgaa ccagctggcg gttttaggac tgccggctgt ggacgccgcc
                                                                      2280
gtagcaggtg aggtctttgc gggccctgcc ctgtcctggt gtctgggact cccgactgtc
                                                                      2340
agtatgatac tgggtctagc aaacctggtg ttgtacttta gatggctggg accccaacgc
                                                                      2400
cttatgttcc ttgtgttgtg gaagctcgct cggggagctt tcccgctggc acttctgatg
                                                                      2460
gggatttegg egactegegg gegeacetee gtgetegggg eegagttetg ettegatget
                                                                      2520
acattcgagg tggacacttc ggtgttgggc tgggtggtgg ccagcgtggt ggcttgggcc
                                                                      2580
atagcgctcc tgagctcgat gagcgcaggg gggtggaggc acaaagccgt gatctatagg
                                                                      2640
acgtggtgta aagggtacca ggctatacgc cagagagtgg ttcggagccc cttcggggag
                                                                      2700
gggcgaccta ctaagcctct gactttcgct tggtgcttgg cctcgtacat ctggccggat
                                                                      2760
gctgtgatga tggtggtggt cgccttggtc ttactctttg gcctgttcga tgcgttggac
                                                                      2820
tgggctttgg aggagatett ggtgteeegg eeetegttge geegtttgge tegggtgate
                                                                      2880
gagtgctgtg tgatggcggg tgagaaggcc acaactgtcc ggctggtctc caagatgtgc
                                                                      2940
gcgagagggg cctacttgtt cgatcatatg ggctcatttt cgcgcgctgt caaggagcgc
                                                                      3000
ctgttggagt gggacgcggc tcttgaacct ctgtcattca ctaqqacqqa ctqtcqcatt
                                                                      3060
ataagagatg ccgctaggac tttgtcttgc ggacagtgcg tcatgggctt gcccgtggta
                                                                      3120
gegegeegtg gagatgaggt teteategge gtettecagg atgtgaatea tttgeeteet
                                                                      3180
gggtttgtcc caaccgcacc agttgtcatc cggcggtgcg gaaagggttt cctaggggtc
                                                                      3240
acaaaggctg ccttgacggg tcgggatcct gacttacatc caggaaacgt catggtgttg
                                                                      3300
gggacggcaa cgtcgcgaag catgggaaca tgtctgaacg gcctgttgtt cacgaccttc
                                                                      3360
catggggctt catcccgaac catcgccacg cccgtggggg cccttaaccc taggtggtgg
                                                                      3420
teegecagtg atgacgteac ggtgtateeg etteeagatg gggcgaette gttgacacet
                                                                      3480
tgcacttgcc aagctgagtc ctgttgggtc atcagatctg acggggcttt atgccatggc
                                                                      3540
ttgagcaagg gggacaaggt agagctggat gtagctatgg aggtctctga cttccgtggt
                                                                      3600
tegteegget egeeggteet gtgegaegaa gggeaegeag taggaatget egtgteagtg
                                                                      3660
ctccattcgg cgggagaagt caccgcggct cgatttacta gggcgtggac ccaagttcca
                                                                      3720
actgacgcca aaaccaccac tgacccccct ccggtgccgg caaaaggagt tttcaaagag
                                                                      3780
gccccgttgt ttatgcctac gggagcggga aagagcactc gcgtcccgtt ggagtacggc
                                                                      3840
aacatggggc acaaggtett gattttgaac ceeteggteg etaetgtgeg ggecatggge
                                                                      3900
ccctacatgg agcggctggc gggcaaacat ccaagcattt actgtggaca tgacaccact
                                                                      3960
gctttcacga ggatcactga ctcccccta acgtattcta cctatgggag gtttttagcc
                                                                      4020
aaccctaggc agatgctacg gggtgtatcg gtggtcatct gtgatgagtg ccacagccat
                                                                      4080
gactcaaccg tgctgctggg cattgggagg gtccgggagc tggcgcgtgg gtgtggagtg
                                                                      4140
caactggtgc tctacgctac cgccacgcct cctggttccc ccatgacaca gcacccttct
                                                                      4200
ataattgaga caaagctgga tgtgggtgag atcccctttt atgggcatgg catacccctc
                                                                      4260
gaacggatgc gaaccgggag gcatctcgta ttctqccatt ccaaqqctqa qtqtqaqcqc
                                                                      4320
cttgccggcc agttctccgc gaggggggta aatgctattg cttattatag ggggaaagac
                                                                      4380
agttecatea tteaggaegg agaeetagtg gtgtgegeea etgaegeaet ttecaeeggg
                                                                      4440
tatactggga attttgattc tgtcaccgat tgtgggttgg tggtggagga ggtcgttgag
                                                                      4500
gtgaccettg atcccaccat taccatetee etgegaacag tgcetgette ggetgaattg
                                                                      4560
tegatgeaga gacgaggaeg caegggtagg ggeaggtetg ggegetaeta etaegegggg
                                                                      4620
gtgggcaaag cccccgctgg ggtggtgcgc tcaggtcctg tctggtcggc ggtggaagcc
                                                                      4680
ggagtgacct ggtacggaat ggaacctgac ctgacagcta acctattgag actttacgac
                                                                      4740
gactgeeett acaccgeage egtegeaget gacattggag aggeegeggt gttettetee
                                                                      4800
gggetegece egitgaggat geacecegat gitagetggg caaaagtieg eggegteaac
                                                                      4860
tggcccctct tggtgggtgt tcagcggacc atgtgccggg aaacactgtc tcctggtcca
                                                                      4920
teggatgace eccaatggge aggtetgaag ggeeegaate etgteeeaet aetgetgagg
                                                                      4980
tggggcaatg atttaccatc taaagtggcc ggccatcaca tagtggacga cctggtccgt
                                                                      5040
agacteggtg tggeggaggg ttatgteege tgegatgegg ggeegatett gatggteggt
                                                                      5100
```

ctcgctatcg cggggggat gatctacgcg tcatacaccg ggtccatagt ggtggtgaca 5160 gactgggatg tgaagggggg tggcgcccc ctttatcggc atggagacca ggccacgcca 5220 caaccggtgg tgcaggtccc cccggtagac catcggccgg ggggggagtc tgcgccatcg 5280 5340 gacgccaaga cagtgacaga tgcggtggca gccatccagg tggattgcga ttggtcagtt atgactctgt cgatcggaga agtactgtcc ttggctcagg ctaagacggc cgaggcctac 5400 acagcaaccg ccaagtggct tgctggctgc tatacgggga cgcgggccgt ccccactgtt 5460 5520 tcaattgttg acaagctctt cgccggaggg tgggcggccg tggtgggcca ttgccacagc 5580 gtaatagctg cggcggtggc ggcctacggg gcttctagga gcccgccgtt ggcagccgcg 5640 getteetace tgatggggtt gggegtegga ggeaaegeee aaaegegttt ggeatetgee 5700 ctectattgg gggetgetgg gacegeeetg ggtacteegg ttgtgggttt gaceatgget ggggcattca tggggggtgc tagcgtctcc ccctctctgg tcaccatcct gttgggggcc 5760 gtgggaggct gggagggcgt tgtcaatgcg gcaagccttg tctttgactt catggcgggg 5820 aaactttcat cagaagattt gtggtatget atcccggtgc tgaccagtcc gggggcgggc 5880 ctcgcgggga tcgcccttgg actggttctg tactcagcta acaactctgg cactaccact 5940 tggctgaacc gtctgctgac gacgttgcca aggtcttcat gcatccctga cagttacttt 6000 cagcaggccg actactgtga caaagtetca getgtgetee geegtttgag cettaecege 6060 actgtggttg ccctggtcaa cagggagccc aaggtggatg aggttcaggt ggggtacgtc 6120 tgggacctgt gggagtggat catgcgccag gtgcgcatgg tcatggctag actcagggcc 6180 ctctgccccg tggtgtcgct gcctttgtgg cactgcgggg aggggtggtc tggggaatgg 6240 ctgttggacg gccatgttga gagtcgctgc ctttgtggct gcgtgatcac cggcgatgtt 6300 ctgaatgggc aactcaaaga accagtttac tctaccaagc tgtgccggca ctattggatg 6360 gggactgttc ctgtgaacat gctgggctat ggtgagacgt cgcctctctt ggcttctgac 6420 accocgaagg tggtaccttt cgggacgtct ggctgggctg aggtggtggt gacccctacc 6480 cacgtagtga tcaggaggac ctccgcctat aagttgctgc gccagcaaat cctatcggct 6540 gctgtagctg agccctatta cgtcgacggc attccggtct cgtgggacgc ggacgcgcga 6600 6660 gegecegeca tggtetatgg ecetgggeaa agtgttacea ttgaegggga gegetaeaee ttgcctcacc agttgaggct taggaacgtg gcgccctctg aggtttcatc cgagatgacc 6720 attgacettg ggacggagac tgaagattca gaactgactg aagccgatet geegeeggeg 6780 gctgcagccc tccaggcgat cgagaatgct gcgaggattc tcgagccgct cattgatgtc 6840 atcatggagg actgtagtac accetetett tgtggtagta geegagagat geetgtgtgg 6900 ggaggagaca teccegcae tecategeca geaettatet eggttaetga gageagetea 6960 gatgagaaga ccccgtcggt gtcctcttcg caggaggata ccccgtcctc agactcattc 7020 gaggtcatcc aagagtccga gacagccgag ggggaggaaa gtgtcttcaa cgtggctctt 7080 tccgtactaa aagccttatt tccacagagc gacgcgacca ggaagctcac cgtcaagatg 7140 tcatgctgtg ctgaaaagag cgtcacacgt tttttttcat tagggctgac ggtggctgat 7200 gttgctagcc tgtgtgagat ggagattcag aaccatacag cctattgtga caaggtgcgc 7260 actccgcttg aattgcaggt tgggtgcttg gtgggcaatg aacttacctt tgaatgtgac 7320 aagtgtgagg ctaggcaaga gactttggcc teetteteet acatetggte tggtgtgeeg 7380 ctaacgcggg ccacgccggc caagccccct gtggtgaggc cggttggttc tttgttggtg 7440 gccgacacta ccaaagtgta tgttaccaat ccggacaatg tgggaaggag ggtggacaag 7500 gtgacettet ggegegetee tagggtteat gacaaatate ttgtggacte tategagegg 7560 gctaagaggg ctgctcaagc ctgcctaagc atgggttaca cttatgagga agcaataagg 7620 7680 actgtaaggc cacatgctgc catgggctgg ggatctaagg tgtcggtcaa agacttggcc accectgegg ggaagatgge tgtecaegat egactteagg agataettga agggaeteeg 7740 gtccccttta cccttactgt gaaaaaggag gtgttcttca aagaccgaaa ggaggagaag 7800 gcccctcgcc tcattgtgtt ccccccctg gacttccgga tagctgaaaa gcttatcctg 7860 ggagaccegg ggegggtage caaggeggtg ttgggggggg ettaegeett ceagtacace 7920 ccaaaccagc gagttaagga gatgctcaaa ctgtgggagt ccaaaaaaaac accttgcgcc 7980 atctgcgtgg acgccacgtg cttcgacagt agcattactg aagaggacgt ggccttagag 8040 acagagetat atgetetgge etetgaceat ceagaatggg tgegageeet tgggaaatae 8100 tatgetteag geaceatggt caceceagaa ggggtgeeeg ttggtgagag gtattgtaga 8160 tecteagggg tettgacaac cagtgegage aattgettga catgetacat caaggtggaa 8220 gccgcctgtg agagggtggg tctgaaaaat gtctcactcc tcattgctgg cgatgactgt 8280 ttgatcatat gtgageggee agtttgegae cetagegaeg etetgggeag ageeetgget 8340 agctatgggt acgcatgcga gccctcatat catgcatcat tggacacggc ccccttctgc 8400 tecaettgge ttgeegagtg taatgeagat gggaagegee atttetteet gaeaaeggae 8460 ttccggaggg cgctcgctcg catgtcgagc gagtatagtg acccgatggc ttcggccatc 8520 ggttacatec teetttatee ttggeatece ateacaeggt gggteateat ecceeaegtg 8580 ctaacatgcg cgtttagggg tggtggtaca ccgtctgatc cggtttggtg ccaagtacat 8640 ggtaattact acaaqtttcc actqqacaaa ctqcctaaca tcatcqtqqc cctccacqqa 8700 ccagcagcgt tgagggttac cgcagacaca actaagacaa aaatggaggc cggtaaggtc 8760 ttaagegace teaageteee tggettageg gteeaeegta agaaggeegg ggeaetgegt 8820

```
acgcgcatgc tccgttcgcg cggttgggcc gagttggcta ggggcctgtt gtggcatcca
                                                                      8880
ggcctacggc tccctcccc tgagattgct ggtatccccg ggggtttccc ctctccccc
                                                                      8940
ccctttatgg gggtggttca tcaattggat ttcacaagcc agaggagtcg ctggcggggg
                                                                      9000
ttgggggtct tagccctgct catcgtggcc ctcttcgggt gaactaaatt catctgttgc
                                                                      9060
ggcaaggtct ggtgactgat catcaccgga ggaggttccc gccctccccg ccccaggggt
                                                                      9120
eteccegetg ggtaaaaagg geeggeett gggaggeatg gtggttaeta acceeetgge
                                                                      9180
agggtcaaag cctgatggtg ctaatgcact gccacttcgg tggcgggtcg ctaccttata
                                                                      9240
gcgtaatccg tgactacggg ctgctcgcag agccctcccc ggatggggca cagtgcactg
                                                                      9300
tgatctgaa
                                                                      9309
```

<210> 20 <211> 2842 <212> PRT <213> Hepatitis G virus

<400> 20 Met Ala Val Leu Leu Leu Leu Val Val Glu Ala Gly Ala Ile Leu 1 Ala Pro Ala Thr His Ala Cys Arg Ala Asn Gly Gln Tyr Phe Leu Thr Asn Cys Cys Ala Pro Glu Asp Ile Gly Phe Cys Leu Glu Gly Gly Cys Leu Val Ala Leu Gly Cys Thr Val Cys Thr Asp Gln Cys Trp Pro Leu Tyr Gln Ala Gly Leu Ala Val Arg Pro Gly Lys Ser Ala Ala Gln Leu 65 70 75 Val Gly Glu Leu Gly Ser Leu Tyr Gly Pro Leu Ser Val Ser Ala Tyr 90 Val Ala Gly Ile Leu Gly Leu Gly Glu Val Tyr Ser Gly Val Leu Thr 105 110 Val Gly Val Ala Leu Thr Arg Arg Val Tyr Pro Val Pro Asn Leu Thr 120 Cys Ala Val Glu Cys Glu Leu Lys Trp Glu Ser Glu Phe Trp Arg Trp 130 135 Thr Glu Gln Leu Ala Ser Asn Tyr Trp Ile Leu Glu Tyr Leu Trp Lys 150 155 160 Val Pro Phe Asp Phe Trp Arg Gly Val Met Ser Leu Thr Pro Leu Leu 170 175 Val Cys Val Ala Ala Leu Leu Leu Glu Gln Arg Ile Val Met Val 185 190 Phe Leu Leu Val Thr Met Ala Gly Met Ser Gln Gly Ala Pro Ala Ser 200 Val Leu Gly Ser Arg Pro Phe Asp Tyr Gly Leu Thr Trp Gln Thr Cys 215 220 Ser Cys Lys Ala Asn Gly Ser Arg Ile Pro Thr Gly Glu Lys Val Trp 230 Asp Arg Gly Asn Val Thr Leu Leu Cys Asp Cys Pro Asn Gly Pro Trp 250 Val Trp Leu Pro Ala Phe Cys Gln Ala Val Gly Trp Gly Asp Pro Ile 265 270 Thr His Trp Ser His Gly Gln Asn Arg Trp Pro Leu Ser Cys Pro Gln 275 280 Tyr Val Tyr Gly Ser Val Ser Val Thr Cys Val Trp Gly Ser Ala Ser 295 300 Trp Phe Ala Ser Thr Gly Gly Arg Asp Ser Lys Ile Asp Val Trp Ser 310 315 Leu Val Pro Val Gly Ser Ala Thr Cys Thr Ile Ala Ala Leu Gly Ser 325 330 Ser Asp Arg Asp Thr Val Val Glu Leu Ser Glu Trp Gly Val Pro Cys 340 345

```
Val Thr Cys Ile Leu Asp Arg Arg Pro Ala Ser Cys Gly Thr Cys Val
                           360
       355
Arg Asp Cys Trp Pro Glu Thr Gly Ser Val Arg Phe Pro Phe His Arg
                        375
                                            380
Cys Gly Thr Gly Pro Arg Leu Thr Lys Asp Leu Glu Ala Val Pro Phe
                   390
                                        395
Val Asn Arg Thr Thr Pro Phe Thr Ile Arg Gly Pro Leu Gly Asn Gln
               405
                                    410
Gly Arg Gly Asn Pro Val Arg Ser Pro Leu Gly Phe Gly Ser Tyr Thr
                               425
Met Thr Lys Ile Arg Asp Thr Leu His Leu Val Lys Cys Pro Thr Pro
        435
                            440
                                                445
Ala Ile Glu Pro Pro Thr Gly Thr Phe Gly Phe Phe Pro Gly Thr Pro
                        455
                                            460
Pro Leu Asn Asn Cys Met Leu Leu Gly Thr Glu Val Ser Glu Ala Leu
                    470
                                        475
Gly Gly Ala Gly Leu Thr Gly Gly Phe Tyr Glu Pro Leu Val Arg Arg
                                    490
Cys Ser Glu Leu Met Gly Arg Arg Asn Pro Val Cys Pro Gly Phe Ala
                                505
                                                    510
Trp Leu Ser Ser Gly Arg Pro Asp Gly Phe Ile His Val Gln Gly His
                            520
Leu Gln Glu Val Asp Ala Gly Asn Phe Ile Pro Pro Pro Arg Trp Leu
                        535
                                            540
Leu Leu Asp Phe Val Phe Val Leu Leu Tyr Leu Met Lys Leu Ala Glu
                                        555
Ala Arg Leu Val Pro Leu Ile Leu Leu Leu Leu Trp Trp Val Asn
                                    570
Gln Leu Ala Val Leu Gly Leu Pro Ala Val Asp Ala Ala Val Ala Gly
                                585
Glu Val Phe Ala Gly Pro Ala Leu Ser Trp Cys Leu Gly Leu Pro Thr
                            600
Val Ser Met Ile Leu Gly Leu Ala Asn Leu Val Leu Tyr Phe Arg Trp
                        615
Leu Gly Pro Gln Arg Leu Met Phe Leu Val Leu Trp Lys Leu Ala Arg
                    630
                                        635
Gly Ala Phe Pro Leu Ala Leu Leu Met Gly Ile Ser Ala Thr Arq Gly
                645
                                    650
                                                        655
Arg Thr Ser Val Leu Gly Ala Glu Phe Cys Phe Asp Ala Thr Phe Glu
                                665
                                                    670
Val Asp Thr Ser Val Leu Gly Trp Val Val Ala Ser Val Val Ala Trp
        675
                            680
Ala Ile Ala Leu Leu Ser Ser Met Ser Ala Gly Gly Trp Arg His Lys
                       695
                                            700
Ala Val Ile Tyr Arg Thr Trp Cys Lys Gly Tyr Gln Ala Ile Arg Gln
                                        715
                    710
Arg Val Val Arg Ser Pro Phe Gly Glu Gly Arg Pro Thr Lys Pro Leu
               725
                                    730
Thr Phe Ala Trp Cys Leu Ala Ser Tyr Ile Trp Pro Asp Ala Val Met
                                745
Met Val Val Ala Leu Val Leu Leu Phe Gly Leu Phe Asp Ala Leu
                            760
Asp Trp Ala Leu Glu Glu Ile Leu Val Ser Arg Pro Ser Leu Arg Arg
                       775
Leu Ala Arg Val Ile Glu Cys Cys Val Met Ala Gly Glu Lys Ala Thr
                   790
                                        795
Thr Val Arg Leu Val Ser Lys Met Cys Ala Arg Gly Ala Tyr Leu Phe
                                   810
Asp His Met Gly Ser Phe Ser Arq Ala Val Lys Glu Arq Leu Leu Glu
           820
                                825
Trp Asp Ala Ala Leu Glu Pro Leu Ser Phe Thr Arg Thr Asp Cys Arg
       835
```

```
Ile Ile Arg Asp Ala Ala Arg Thr Leu Ser Cys Gly Gln Cys Val Met
                     855
                                         860
Gly Leu Pro Val Val Ala Arg Arg Gly Asp Glu Val Leu Ile Gly Val
                                     875
                  870
Phe Gln Asp Val Asn His Leu Pro Pro Gly Phe Val Pro Thr Ala Pro
              885
                                 890
Val Val Ile Arg Arg Cys Gly Lys Gly Phe Leu Gly Val Thr Lys Ala
                              905
Ala Leu Thr Gly Arg Asp Pro Asp Leu His Pro Gly Asn Val Met Val
                         920
Leu Gly Thr Ala Thr Ser Arg Ser Met Gly Thr Cys Leu Asn Gly Leu
                      935
                                         940
Leu Phe Thr Thr Phe His Gly Ala Ser Ser Arg Thr Ile Ala Thr Pro
                  950
                                     955
Val Gly Ala Leu Asn Pro Arg Trp Trp Ser Ala Ser Asp Asp Val Thr
               965
                                 970
Val Tyr Pro Leu Pro Asp Gly Ala Thr Ser Leu Thr Pro Cys Thr Cys
                              985
                                                990
Gln Ala Glu Ser Cys Trp Val Ile Arg Ser Asp Gly Ala Leu Cys His
                          1000
                                             1005
Gly Leu Ser Lys Gly Asp Lys Val Glu Leu Asp Val Ala Met Glu Val
                      1015
                                         1020
Ser Asp Phe Arg Gly Ser Ser Gly Ser Pro Val Leu Cys Asp Glu Gly
                  1030
                                     1035
His Ala Val Gly Met Leu Val Ser Val Leu His Ser Ala Gly Glu Val
               1045
                                 1050
Thr Ala Ala Arg Phe Thr Arg Ala Trp Thr Gln Val Pro Thr Asp Ala
           1060
                              1065
Lys Thr Thr Asp Pro Pro Pro Val Pro Ala Lys Gly Val Phe Lys
       1075
                          1080
                                             1085
Glu Ala Pro Leu Phe Met Pro Thr Gly Ala Gly Lys Ser Thr Arg Val
                      1095
                                         1100
Pro Leu Glu Tyr Gly Asn Met Gly His Lys Val Leu Ile Leu Asn Pro
                  1110
                                     1115
Ser Val Ala Thr Val Arg Ala Met Gly Pro Tyr Met Glu Arg Leu Ala
               1125
                                 1130
Gly Lys His Pro Ser Ile Tyr Cys Gly His Asp Thr Thr Ala Phe Thr
           1140
                              1145
                                                1150
Arg Ile Thr Asp Ser Pro Leu Thr Tyr Ser Thr Tyr Gly Arg Phe Leu
                          1160
       1155
                                             1165
Ala Asn Pro Arg Gln Met Leu Arg Gly Val Ser Val Val Ile Cys Asp
                      1175
                                         1180
Glu Cys His Ser His Asp Ser Thr Val Leu Leu Gly Ile Gly Arg Val
       1190
                                     1195 .
Arg Glu Leu Ala Arg Gly Cys Gly Val Gln Leu Val Leu Tyr Ala Thr
              1205
                                 1210
Ala Thr Pro Pro Gly Ser Pro Met Thr Gln His Pro Ser Ile Ile Glu
                              1225
                                                1230
Thr Lys Leu Asp Val Gly Glu Ile Pro Phe Tyr Gly His Gly Ile Pro
                          1240
                                             1245
Leu Glu Arg Met Arg Thr Gly Arg His Leu Val Phe Cys His Ser Lys
                      1255
                                         1260
Ala Glu Cys Glu Arg Leu Ala Gly Gln Phe Ser Ala Arg Gly Val Asn
                 1270
                                     1275
Ala Ile Ala Tyr Tyr Arg Gly Lys Asp Ser Ser Ile Ile Gln Asp Gly
                                1290
              1285
Asp Leu Val Val Cys Ala Thr Asp Ala Leu Ser Thr Gly Tyr Thr Gly
          1300
                            1305
Asn Phe Asp Ser Val Thr Asp Cys Gly Leu Val Val Glu Glu Val Val
                          1320
Glu Val Thr Leu Asp Pro Thr Ile Thr Ile Ser Leu Arg Thr Val Pro
   1330
                      1335
```

```
Ala Ser Ala Glu Leu Ser Met Gln Arg Arg Gly Arg Thr Gly Arg Gly
                                     1355
            1350
Arg Ser Gly Arg Tyr Tyr Ala Gly Val Gly Lys Ala Pro Ala Gly
               1365
                                 1370
Val Val Arg Ser Gly Pro Val Trp Ser Ala Val Glu Ala Gly Val Thr
           1380
                             1385
                                                1390
Trp Tyr Gly Met Glu Pro Asp Leu Thr Ala Asn Leu Leu Arg Leu Tyr
       1395
                          1400
                                          1405
Asp Asp Cys Pro Tyr Thr Ala Ala Val Ala Ala Asp Ile Gly Glu Ala
                      1415
                                         1420
Ala Val Phe Phe Ser Gly Leu Ala Pro Leu Arg Met His Pro Asp Val
                  1430
                         1435
Ser Trp Ala Lys Val Arg Gly Val Asn Trp Pro Leu Leu Val Gly Val
               1445
                                 1450
Gln Arg Thr Met Cys Arg Glu Thr Leu Ser Pro Gly Pro Ser Asp Asp
           1460
                              1465
                                                1470
Pro Gln Trp Ala Gly Leu Lys Gly Pro Asn Pro Val Pro Leu Leu
       1475
                          1480
                                             1485
Arg Trp Gly Asn Asp Leu Pro Ser Lys Val Ala Gly His His Ile Val
                      1495
                                         1500
Asp Asp Leu Val Arg Arg Leu Gly Val Ala Glu Gly Tyr Val Arg Cys
                  1510
                                     1515
Asp Ala Gly Pro Ile Leu Met Val Gly Leu Ala Ile Ala Gly Gly Met
               1525
                                 1530
                                                    1535
Ile Tyr Ala Ser Tyr Thr Gly Ser Ile Val Val Val Thr Asp Trp Asp
           1540
                              1545
                                                1550
Val Lys Gly Gly Gly Ala Pro Leu Tyr Arg His Gly Asp Gln Ala Thr
       1555
                          1560
                                             1565
Pro Gln Pro Val Val Gln Val Pro Pro Val Asp His Arg Pro Gly Gly
                      1575
                                         1580
Glu Ser Ala Pro Ser Asp Ala Lys Thr Val Thr Asp Ala Val Ala Ala
                  1590
                                     1595
Ile Gln Val Asp Cys Asp Trp Ser Val Met Thr Leu Ser Ile Gly Glu
               1605
                                 1610
Val Leu Ser Leu Ala Gln Ala Lys Thr Ala Glu Ala Tyr Thr Ala Thr
           1620
                              1625
Ala Lys Trp Leu Ala Gly Cys Tyr Thr Gly Thr Arg Ala Val Pro Thr
                          1640
                                            1645
Val Ser Ile Val Asp Lys Leu Phe Ala Gly Gly Trp Ala Ala Val Val
                      1655
                                         1660
Gly His Cys His Ser Val Ile Ala Ala Ala Val Ala Ala Tyr Gly Ala
                  1670
                                     1675
Ser Arg Ser Pro Pro Leu Ala Ala Ala Ser Tyr Leu Met Gly Leu
                                 1690
Gly Val Gly Gly Asn Ala Gln Thr Arg Leu Ala Ser Ala Leu Leu
           1700
                             1705
                                                1710
Gly Ala Ala Gly Thr Ala Leu Gly Thr Pro Val Val Gly Leu Thr Met
                          1720
                                            1725
Ala Gly Ala Phe Met Gly Gly Ala Ser Val Ser Pro Ser Leu Val Thr
                      1735
                                         1740
Ile Leu Leu Gly Ala Val Gly Gly Trp Glu Gly Val Val Asn Ala Ala
                  1750
                                     1755
Ser Leu Val Phe Asp Phe Met Ala Gly Lys Leu Ser Ser Glu Asp Leu
              1765
                                 1770
Trp Tyr Ala Ile Pro Val Leu Thr Ser Pro Gly Ala Gly Leu Ala Gly
          1780
                             1785
                                                1790
Ile Ala Leu Gly Leu Val Leu Tyr Ser Ala Asn Asn Ser Gly Thr Thr
       1795
                         1800
Thr Trp Leu Asn Arg Leu Leu Thr Thr Leu Pro Arg Ser Ser Cys Ile
                     1815
                                        1820
Pro Asp Ser Tyr Phe Gln Gln Ala Asp Tyr Cys Asp Lys Val Ser Ala
                  1830
                                     1835
```

```
Val Leu Arg Arg Leu Ser Leu Thr Arg Thr Val Val Ala Leu Val Asn
                              1850 1855
             1845
Arg Glu Pro Lys Val Asp Glu Val Gln Val Gly Tyr Val Trp Asp Leu
                           1865
         1860
                                            1870
Trp Glu Trp Ile Met Arg Gln Val Arg Met Val Met Ala Arg Leu Arg
                    1880
       1875
                                         1885
Ala Leu Cys Pro Val Val Ser Leu Pro Leu Trp His Cys Gly Glu Gly
                                     1900
           1895
Trp Ser Gly Glu Trp Leu Leu Asp Gly His Val Glu Ser Arg Cys Leu
        1910
                              1915 1920
Cys Gly Cys Val Ile Thr Gly Asp Val Leu Asn Gly Gln Leu Lys Glu
             1925 1930
Pro Val Tyr Ser Thr Lys Leu Cys Arg His Tyr Trp Met Gly Thr Val
                         1945
          1940
                                            1950
Pro Val Asn Met Leu Gly Tyr Gly Glu Thr Ser Pro Leu Leu Ala Ser
       1955
                       1960
                                         1965
Asp Thr Pro Lys Val Val Pro Phe Gly Thr Ser Gly Trp Ala Glu Val
                    1975
                                      1980
Val Val Thr Pro Thr His Val Val Ile Arg Arg Thr Ser Ala Tyr Lys
                 1990
                                  1995
Leu Leu Arg Gln Gln Ile Leu Ser Ala Ala Val Ala Glu Pro Tyr Tyr
             2005
                               2010
Val Asp Gly Ile Pro Val Ser Trp Asp Ala Asp Ala Arg Ala Pro Ala
          2020
                          2025
Met Val Tyr Gly Pro Gly Gln Ser Val Thr Ile Asp Gly Glu Arg Tyr
       2035
                        2040
                                         2045
Thr Leu Pro His Gln Leu Arg Leu Arg Asn Val Ala Pro Ser Glu Val
                    2055
Ser Ser Glu Met Thr Ile Asp Leu Gly Thr Glu Thr Glu Asp Ser Glu
                 2070
                                  2075
Leu Thr Glu Ala Asp Leu Pro Pro Ala Ala Ala Leu Gln Ala Ile
             2085
                               2090
Glu Asn Ala Ala Arg Ile Leu Glu Pro Leu Ile Asp Val Ile Met Glu
          2100
                           2105
Asp Cys Ser Thr Pro Ser Leu Cys Gly Ser Ser Arg Glu Met Pro Val
      2115 2120
                                         2125
Trp Gly Gly Asp Ile Pro Arg Thr Pro Ser Pro Ala Leu Ile Ser Val
                    2135
                                      2140
Thr Glu Ser Ser Ser Asp Glu Lys Thr Pro Ser Val Ser Ser Ser Gln
                 2150
                                  2155
Glu Asp Thr Pro Ser Ser Asp Ser Phe Glu Val Ile Gln Glu Ser Glu
             2165
                               2170
Thr Ala Glu Gly Glu Glu Ser Val Phe Asn Val Ala Leu Ser Val Leu
                           2185
Lys Ala Leu Phe Pro Gln Ser Asp Ala Thr Arg Lys Leu Thr Val Lys
                       2200
                                         2205
Met Ser Cys Cys Ala Glu Lys Ser Val Thr Arg Phe Phe Ser Leu Gly
                    2215
                                     2220
Leu Thr Val Ala Asp Val Ala Ser Leu Cys Glu Met Glu Ile Gln Asn
                2230
                                  2235
His Thr Ala Tyr Cys Asp Lys Val Arg Thr Pro Leu Glu Leu Gln Val
             2245
                               2250
Gly Cys Leu Val Gly Asn Glu Leu Thr Phe Glu Cys Asp Lys Cys Glu
          2260
                           2265
Ala Arg Gln Glu Thr Leu Ala Ser Phe Ser Tyr Ile Trp Ser Gly Val
      2275
                       2280
Pro Leu Thr Arg Ala Thr Pro Ala Lys Pro Pro Val Val Arg Pro Val
                   2295
                                     2300
Gly Ser Leu Leu Val Ala Asp Thr Thr Lys Val Tyr Val Thr Asn Pro
                2310
                                  2315
Asp Asn Val Gly Arg Arg Val Asp Lys Val Thr Phe Trp Arg Ala Pro
             2325
                              2330
```

Arg Val His Asp Lys Tyr Leu Val Asp Ser Ile Glu Arg Ala Lys Arg Ala Ala Gln Ala Cys Leu Ser Met Gly Tyr Thr Tyr Glu Glu Ala Ile Arg Thr Val Arg Pro His Ala Ala Met Gly Trp Gly Ser Lys Val Ser Val Lys Asp Leu Ala Thr Pro Ala Gly Lys Met Ala Val His Asp Arg Leu Gln Glu Ile Leu Glu Gly Thr Pro Val Pro Phe Thr Leu Thr Val Lys Lys Glu Val Phe Phe Lys Asp Arg Lys Glu Glu Lys Ala Pro Arg Leu Ile Val Phe Pro Pro Leu Asp Phe Arg Ile Ala Glu Lys Leu Ile Leu Gly Asp Pro Gly Arg Val Ala Lys Ala Val Leu Gly Gly Ala Tyr Ala Phe Gln Tyr Thr Pro Asn Gln Arg Val Lys Glu Met Leu Lys Leu Trp Glu Ser Lys Lys Thr Pro Cys Ala Ile Cys Val Asp Ala Thr Cys Phe Asp Ser Ser Ile Thr Glu Glu Asp Val Ala Leu Glu Thr Glu Leu Tyr Ala Leu Ala Ser Asp His Pro Glu Trp Val Arg Ala Leu Gly Lys Tyr Tyr Ala Ser Gly Thr Met Val Thr Pro Glu Gly Val Pro Val Gly Glu Arg Tyr Cys Arg Ser Ser Gly Val Leu Thr Thr Ser Ala Ser Asn Cys Leu Thr Cys Tyr Ile Lys Val Glu Ala Ala Cys Glu Arg Val Gly Leu Lys Asn Val Ser Leu Leu Ile Ala Gly Asp Asp Cys Leu Ile Ile Cys Glu Arg Pro Val Cys Asp Pro Ser Asp Ala Leu Gly Arg Ala Leu Ala Ser Tyr Gly Tyr Ala Cys Glu Pro Ser Tyr His Ala Ser Leu Asp Thr Ala Pro Phe Cys Ser Thr Trp Leu Ala Glu Cys Asn Ala Asp Gly Lys Arg His Phe Phe Leu Thr Thr Asp Phe Arg Arg Ala Leu Ala Arg Met Ser Ser Glu Tyr Ser Asp Pro Met Ala Ser Ala Ile Gly Tyr Ile Leu Leu Tyr Pro Trp His Pro Ile Thr Arg Trp Val Ile Ile Pro His Val Leu Thr Cys Ala Phe Arg Gly Gly Gly Thr Pro Ser Asp Pro Val Trp Cys Gln Val His Gly Asn Tyr Tyr Lys Phe Pro Leu Asp Lys Leu Pro Asn Ile Ile Val Ala Leu His Gly Pro Ala Ala Leu Arg Val Thr Ala Asp Thr Thr Lys Thr Lys Met Glu Ala Gly Lys Val Leu Ser Asp Leu Lys Leu Pro Gly Leu Ala Val His Arg Lys Lys Ala Gly Ala Leu Arg Thr Arg Met Leu Arg Ser Arg Gly Trp Ala Glu Leu Ala Arg Gly Leu Leu Trp His Pro Gly Leu Arg Leu Pro Pro Pro Glu Ile Ala Gly Ile Pro Gly Gly Phe Pro Ser Pro Pro Pro Phe Met Gly Val Val His Gln Leu Asp Phe Thr Ser Gln Arg Ser Arg Trp Arg Gly Leu Gly Val

Leu Ala Leu Leu Ile Val Ala Leu Phe Gly 2835 2840

<210> 21

<211> 414 <212> PRT <213> Hepatitis G virus Val Ala Leu Val Asn Arg Glu Pro Lys Val Asp Glu Val Gln Val Gly Tyr Val Trp Asp Leu Trp Glu Trp Ile Met Arg Gln Val Arg Met Val Met Ala Arg Leu Arg Ala Leu Cys Pro Val Val Ser Leu Pro Leu Trp His Cys Gly Glu Gly Trp Ser Gly Glu Trp Leu Leu Asp Gly His Val Glu Ser Arg Cys Leu Cys Gly Cys Val Ile Thr Gly Asp Val Leu Asn Gly Gln Leu Lys Glu Pro Val Tyr Ser Thr Lys Leu Cys Arg His Tyr Trp Met Gly Thr Val Pro Val Asn Met Leu Gly Tyr Gly Glu Thr Ser 100 105 Pro Leu Leu Ala Ser Asp Thr Pro Lys Val Val Pro Phe Gly Thr Ser 120 Gly Trp Ala Glu Val Val Thr Pro Thr His Val Val Ile Arg Arg 130 135 Thr Ser Ala Tyr Lys Leu Leu Arg Gln Gln Ile Leu Ser Ala Ala Val 150 Ala Glu Pro Tyr Tyr Val Asp Gly Ile Pro Val Ser Trp Asp Ala Asp 165 170 Ala Arg Ala Pro Ala Met Val Tyr Gly Pro Gly Gln Ser Val Thr Ile 185 Asp Gly Glu Arg Tyr Thr Leu Pro His Gln Leu Arg Leu Arg Asn Val 200 Ala Pro Ser Glu Val Ser Ser Glu Met Thr Ile Asp Leu Gly Thr Glu 215 220 Thr Glu Asp Ser Glu Leu Thr Glu Ala Asp Leu Pro Pro Ala Ala Ala 230 235 Ala Leu Gln Ala Ile Glu Asn Ala Ala Arg Ile Leu Glu Pro Leu Ile 250 Asp Val Ile Met Glu Asp Cys Ser Thr Pro Ser Leu Cys Gly Ser Ser 265 270 Arg Glu Met Pro Val Trp Gly Gly Asp Ile Pro Arg Thr Pro Ser Pro 280 Ala Leu Ile Ser Val Thr Glu Ser Ser Ser Asp Glu Lys Thr Pro Ser 295 300 Val Ser Ser Ser Gln Glu Asp Thr Pro Ser Ser Asp Ser Phe Glu Val 310 315 Ile Gln Glu Ser Glu Thr Ala Glu Gly Glu Glu Ser Val Phe Asn Val 325 330 Ala Leu Ser Val Leu Lys Ala Leu Phe Pro Gln Ser Asp Ala Thr Arg 340 345 Lys Leu Thr Val Lys Met Ser Cys Cys Ala Glu Lys Ser Val Thr Arg 360 Phe Phe Ser Leu Gly Leu Thr Val Ala Asp Val Ala Ser Leu Cys Glu 375 380 Met Glu Ile Gln Asn His Thr Ala Tyr Cys Asp Lys Val Arg Thr Pro 390 395 Leu Glu Leu Gln Val Gly Cys Leu Val Gly Asn Glu Leu Thr 405 410

```
<211> 414
<212> PRT
<213> Hepatitis G virus
<400> 22
Val Ala Leu Val Asn Arg Glu Pro Lys Val Asp Glu Val Gln Val Gly
                                    10
Tyr Val Trp Asp Leu Trp Glu Trp Ile Met Arg Gln Val Arg Met Val
                                25
Met Ala Arg Leu Arg Ala Leu Cys Pro Val Val Ser Leu Pro Leu Trp
                            40
His Cys Gly Glu Gly Trp Ser Gly Glu Trp Leu Leu Asp Gly His Val
Glu Ser Arg Cys Leu Cys Gly Cys Val Ile Thr Gly Asp Val Leu Asn
                    70
Gly Gln Leu Lys Asp Pro Val Tyr Ser Thr Lys Leu Cys Arg His Tyr
                85
Trp Met Gly Thr Val Pro Val Asn Met Leu Gly Tyr Gly Glu Thr Ser
                                105
Pro Leu Leu Ala Ser Asp Thr Pro Lys Val Val Pro Phe Gly Thr Ser
                            120
Gly Trp Ala Glu Val Val Thr Pro Thr His Val Val Ile Arg Arg
                        135
                                            140
Thr Ser Ala Tyr Lys Leu Leu Arg Gln Gln Ile Leu Ser Ala Ala Val
                                        155
Ala Glu Pro Tyr Tyr Val Asp Gly Ile Pro Val Ser Trp Asp Ala Asp
                                    170
Ala Arg Ala Pro Ala Met Val Tyr Gly Pro Gly Gln Ser Val Thr Ile
                                185
Asp Gly Glu Arg Tyr Thr Leu Pro His Gln Leu Arg Leu Arg Asn Val
        195
                            200
Ala Pro Ser Glu Val Ser Ser Glu Val Ser Ile Asp Ile Gly Thr Glu
                        215
                                            220
Thr Glu Asp Ser Glu Leu Thr Glu Ala Asp Leu Pro Pro Ala Ala Ala
                    230
                                        235
Ala Leu Gln Ala Ile Glu Asn Ala Ala Arg Ile Leu Glu Pro His Ile
                245
                                    250
Asp Val Ile Met Glu Asp Cys Ser Thr Pro Ser Leu Cys Gly Ser Ser
                                265
                                                    270
Arg Glu Met Pro Val Trp Gly Glu Asp Ile Pro Arg Thr Pro Ser Pro
                            280
Ala Leu Ile Ser Val Thr Glu Ser Ser Pro Asp Glu Lys Thr Pro Ser
                        295
                                            300
Val Ser Ser Ser Gln Glu Asp Thr Pro Ser Ser Asp Ser Phe Glu Val
                    310
                                        315
Ile Gln Glu Ser Glu Thr Ala Glu Gly Glu Glu Ser Val Phe Asn Val
                                    330
Ala Leu Ser Val Leu Lys Ala Leu Phe Pro Gln Ser Asp Ala Thr Arg
                                345
Lys Leu Thr Val Lys Met Ser Cys Cys Val Glu Lys Ser Val Thr Arg
                            360
Phe Phe Ser Leu Gly Leu Thr Val Ala Asp Val Ala Ser Leu Cys Glu
                        375
Met Glu Ile Gln Asn His Thr Ala Tyr Cys Asp Lys Val Arg Thr Pro
                   390
Leu Glu Leu Gln Val Gly Cys Leu Val Gly Asn Glu Leu Thr
<210> 23
```

<210> 22

<211> 97

<212> DNA

<213> Hepatitis G virus	
	60 97
<210> 24 <211> 97 <212> DNA <213> Hepatitis G virus	
	60 97
<210> 25 <211> 97 <212> DNA <213> Hepatitis G virus	
	60 97
<210> 26 <211> 97 <212> DNA <213> Hepatitis G virus	
	60 97
<210> 27 <211> 97 <212> DNA <213> Hepatitis G virus	
	60 97
<210> 28 <211> 97 <212> DNA <213> Hepatitis G virus	
	60 97
<210> 29 <211> 97 <212> DNA <213> Hepatitis G virus	
	60 97
<210> 30 <211> 97	

<212> DNA <213> Hepatitis G virus	
	60 97
<210> 31 <211> 97 <212> DNA <213> Hepatitis G virus	
	60 97
<210> 32 <211> 97 <212> DNA <213> Hepatitis G virus	
	60 97
<210> 33 <211> 97 <212> DNA <213> Hepatitis G virus	
	60 97
<210> 34 <211> 97 <212> DNA <213> Hepatitis G virus	
	60 97
<210> 35 <211> 97 <212> DNA <213> Hepatitis G virus	
	50 97
<210> 36 <211> 97 <212> DNA <213> Hepatitis G virus	
	50 97

```
<210> 37
<211> 97
<212> DNA
<213> Hepatitis G virus
<400> 37
ccattgacat tgggacggag actgaagact cagaactgac tgaggccgat ctgccgccgg
                                                                          60
cggctgctgc tctccaagcg attgagaatg ctgcgag
                                                                          97
<210> 38
<211> 97
<212> DNA
<213> Hepatitis G virus
<400> 38
ccattgacat tgggacggag actgaagact cagaactgac tgaggccgat ctgccgccgg
                                                                          60
cggctgctgc tctccaagcg attgagaatg ctgcgag
                                                                          97
<210> 39
<211> 97
<212> DNA
<213> Hepatitis G virus
<400> 39
ccattgacat tgggacggag actgaagact cagaactgac tgaggccgat ctgccgccgg
                                                                          60
cggctgctgc tctccaagcg attgagaatg ctgcgag
                                                                          97
<210> 40
<211> 97
<212> DNA
<213> Hepatitis G virus
<400> 40
ccattgacat tgggacggag actgaagact cagaactgac tgaggccgat ctgccgccgg
                                                                         60
cggctgctgc tctccaagcg attgagaatg ctgcgag
                                                                          97
<210> 41
<211> 15
<212> PRT
<213> Hepatitis G virus
<400> 41
Phe Gln Tyr Thr Pro Asn Gln Arg Ile Arg Glu Met Leu Lys Leu
1
                 5
<210> 42
<211> 15
<212> PRT
<213> Hepatitis G virus
<400> 42
Phe Gln Tyr Thr Pro Asn Gln Arg Val Lys Glu Met Leu Lys Leu
<210> 43
<211> 16
<212> PRT
<213> Hepatitis G virus
<400> 43
Gly Ile Pro Gly Ala Phe Pro Leu Ser Pro Pro Tyr Met Gly Val Val
                 5
                                     10
                                                          15
```

. . .

```
<210> 44
<211> 16
<212> PRT
<213> Hepatitis G virus

<400> 44
Gly Ile Pro Gly Gly Phe Ser Pro Ser Pro Pro Phe Met Gly Val Val
1 5 10 15
```